SEQUENCE LISTING

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<110> Mallick, Birendra Nath |
Kar, Santosh Kumar |
Mishra, Bibhuti Bhusan
Madan, Vibha |
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<120> A METHOD FOR IDENTIFICATION AND/OR DIAGNOSIS OF REM SLEEP LOSS FROM BLOOD SAMPLES

<138> JNU-REM

<160> 1

<170> PatentIn version 3.1

<210> 1 | <211> 1477

<212> PRT

<213> Rattus norvegicus

<400> 1

Met Lys Lys Asp Arg Glu Ala Gln Leu Cys Leu Phe Ser Ala Leu Leu 1 5 10 15

Ala Phe Leu Pro Phe Ala Ser Leu Leu Asn Gly Asn Ser Lys Tyr Met 26 25 38

Val Leu Val Pro Ser Gln Leu Tyr Thr Glu Thr Pro Glu Lys Ile Cys 35 40 45

Leu His Leu Tyr His Leu Asn Glu Thr Val Thr Val Thr Ala Ser Leu 50 55 60

Ile Ser Gln Arg Gly Thr Arg Lys Leu Phe Asp Glu Leu Val Val Asp 65 70 75 80

Lys Asp Leu Phe His Cys Val Ser Phe Thr Ile Pro Arg Leu Pro Ser

Ser Glu Glu Glu Glu Ser Leu Asp Ile Asn Ile Glu Gly Ala Lys His 100 105 110

Lys Phe Ser Glu Arg Arg Val Val Leu Val Lys Asn Lys Glu Ser Val 115 120 125

Val Phe Val Gln Thr Asp Lys Pro Met Tyr Lys Pro Gly Gln Ser Val 130 135 140

Lys 145	Phe	Arg	Val	Val	Ser 150	Met	Asp	Lys	Asn	Leu 155	His	Pro	Leu	Asn	G1u 160
Leu	Phe	Pro	Leu	Ala 165	Tyr	Ile	Glu	Asp	Pro 170	Lys	Met	Asn	Arg	Ile 175	Met
Gln	Trp	G1n	Asp 180	Val	Lys	Thr	G1u	Asn 185	G1y	Leu	Lys	G1n	Leu 190	Ser	Phe
Ser	Leu	Ser 195	Ala	G1u	Pro	Ile	61n 200	G1y	Pro	Tyr	Lys	I1e 205	Val	Ile	Leu
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Val 225	Leu	Pro	Arg	Phe	G1y 230	Val	Asp	Val	Lys	Val 235	Pro	Asn	Ala	Ile	Ser 240
Val	Tyr	Asp	G1u	Ile 245	Ile	Asn	Val	Thr	Ala 250	Cys	Ala	Thr	Tyr	Thr 255	Tyr
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Asn	Ala	Thr	Val	Thr 325	Glu	Glu	G1y	Thr	61y 330		Glu	Phe	Ser	61y 335	Ser
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- Lys Ala Arg Asp Ala Gly Tyr Thr Asn Ala Thr Thr Thr Asp Gln His 385 390 395 400
- Gly Leu Ala Lys Phe Ser Ile Asp Thr Asn Gly Ile Ser Asp Tyr Ser 405 410 415
- Leu Asn Ile Lys Val Tyr His Lys Glu Glu Ser Ser Cys Ile His Ser 420 425 430
- Ser Cys Thr Ala Glu Arg His Ala Glu Ala His His Thr Ala Tyr Ala 435 440 445
- Val Tyr Ser Leu Ser Lys Ser Tyr Ile Tyr Leu Asp Thr Glu Ala Gly 450 455 460
- Val Leu Pro Cys Asn Gln Ile His Thr Val Gln Ala His Phe Ile Leu 465 470 475 480
- Lys Gly Gln Val Leu Gly Val Leu Gln Gln Ile Val Phe His Tyr Leu 485 490 495
- Val Met Ala Gln Gly Ser Ile Leu Gln Thr Gly Asn His Thr His Gln 500 505 510
- Val Glu Pro Gly Glu Ser Gln Val Gln Gly Asn Phe Ala Leu Glu Ile 515 520 525
- Pro Val Glu Phe Ser Met Val Pro Val Ala Lys Met Leu Ile Tyr Thr 530 540
- Ile Leu Pro Asp Gly Glu Val Ile Ala Asp Ser Val Lys Phe Gln Val 545 550 555 560
- Glu Lys Cys Leu Arg Asn Lys Val His Leu Ser Phe Ser Pro Ser Gln 565 570 575
- Ser Leu Pro Ala Ser Gln Thr His Met Arg Val Thr Ala Ser Pro Gln 580 585 590
- Ser Leu Cys Gly Leu Arg Ala Val Asp Gln Ser Val Leu Leu Gln Lys 595 600 605

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- Pro Glu Ala Glu Leu Ser Pro Ser Leu Ile Tyr Asp Leu Pro Gly Met 610 615 620
- Gln Asp Ser Asn Phe Ile Ala Ser Ser Asn Asp Pro Phe Glu Asp Glu 625 630 635 640
- Asp Tyr Cys Leu Met Tyr Gln Pro Ile Ala Arg Glu Lys Asp Val Tyr 645 650 655
- Arg Tyr Val Arg Glu Thr Gly Leu Met Ala Phe Thr Asn Leu Lys Ile 669 665 670
- Lys Leu Pro Thr Tyr Cys Asn Thr Asp Tyr Asp Met Val Pro Leu Ala 675 680 685
- Val Pro Ala Val Ala Leu Asp Ser Ser Thr Asp Arg Gly Met Tyr Glu 690 700
- Ser Leu Pro Val Val Ala Val Lys Ser Pro Leu Pro Gln Glu Pro Pro 785 710 715 720
- Arg Lys Asp Pro Pro Pro Lys Asp Pro Val Ile Glu Thr Ile Arg Asn 725 730 735
- Tyr Phe Pro Glu Thr Trp Ile Trp Asp Leu Val Thr Val Asn Ser Ser 748 745 750
- Gly Val Thr Glu Leu Glu Met Thr Val Pro Asp Thr Ile Thr Glu Trp 755 760 765
- Lys Ala Gly Ala Leu Cys Leu Ser Asn Asp Thr Gly Leu Gly Leu Ser 770 775 780
- Ser Val Ala Ser Phe Gln Ala Phe Gln Pro Phe Phe Val Glu Leu Thr 785 790 795 800
- Met Pro Tyr Ser Val Ile Arg Gly Glu Ala Phe Thr Leu Lys Ala Thr 805 810 815
- Val Leu Asn Tyr Leu Pro Thr Ser Leu Pro Met Ala Val Leu Leu Glu 820 825 830
- Ala Ser Pro Asp Phe Thr Ala Val Pro Val Glu Asn Asn Gln Asp Ser 835 840 845

- Tyr Cys Leu Gly Ala Asn Gly Arg His Thr Ser Ser Trp Leu Val Thr 850 855 860
- Pro Lys Ser Leu Gly Asn Val Asn Phe Ser Val Ser Ala Glu Ala Arg 865 870 875 880
- Gln Ser Pro Gly Pro Cys Gly Ser Glu Val Ala Thr Val Pro Glu Thr 885 896 895
- Gly Arg Lys Asp Thr Val Val Lys Val Leu Ile Val Glu Pro Glu Gly 900 905 910
- Ile Lys Lys Glu His Thr Phe Ser Ser Leu Leu Cys Ala Ser Asp Ala 915 920 925
- Glu Leu Ser Glu Thr Leu Ser Leu Leu Leu Pro Pro Thr Val Val Lys 930 935 940
- Asp Ser Ala Arg Ala His Phe Ser Val Met Gly Asp Ile Leu Ser Ser 945 950 955 960
- Ala Ile Lys Asn Thr Gln Asn Leu Ile Gln Met Pro Tyr Gly Cys Gly 965 970 975
- Glu Gln Asn Met Val Leu Phe Ala Pro Asn Ile Tyr Val Leu Lys Tyr 980 998 998
- Leu Asn Glu Thr Gln Gln Leu Thr Glu Lys Ile Lys Ser Lys Ala Leu 995 1000 1005
- Gly Tyr Leu Arg Ala Gly Tyr Gln Arg Glu Leu Asn Tyr Lys His 1010 1015 1020
- Lys Asp Gly Ser Tyr Ser Ala Phe Gly Asp His Asn Gly Gln Gly 1025 1030 1035
- Gln Gly Asn Thr Trp Leu Thr Ala Phe Val Leu Lys Ser Phe Ala 1040 1045 1050
- Gln Ala Arg Ala Phe Ile Phe Ile Asp Glu Ser His Ile Thr Asp 1055 1060 1065

- Ala Phe Thr Trp Leu Ser Lys Gln Gln Lys Asp Ser Gly Cys Phe 1070 1080
- Arg Ser Ser Gly Ser Leu Leu Asn Asn Ala Met Lys Gly Gly Val 1085 1090 1095
- Asp Asp Glu Ile Thr Leu Ser Ala Tyr Ile Thr Met Ala Leu Leu 1100 1105 1110
- Glu Ser Ser Leu Pro Asp Thr Asp Pro Val Val Ser Lys Ala Leu 1115 1120 1125
- Ser Cys Leu Glu Ser Ser Trp Glu Asn Ile Glu Gln Gly Gly Asn 1130 1135 1140
- Gly Ser Phe Val Tyr Thr Lys Ala Leu Met Ala Tyr Ala Phe Ala 1145 1150 1155
- Leu Ala Gly Asn Gln Glu Lys Arg Asn Glu Ile Leu Lys Ser Leu 1160 1165 1170
- Asp Lys Glu Ala Ile Lys Glu Asp Asn Ser Ile His Trp Glu Arg 1175 1180 1185
- Pro Gln Lys Pro Thr Lys Ser Glu Gly Tyr Leu Tyr Thr Pro Gln 1198 1195 1200
- Ala Ser Ser Ala Glu Val Glu Met Ser Ala Tyr Val Val Leu Ala 1205 1210 1215
- Arg Leu Thr Ala Gln Pro Ala Pro Ser Pro Glu Asp Leu Ala Leu 1220 1225 1230
- Ser Met Gly Thr Ile Lys Trp Leu Thr Lys Gln Gln Asn Ser Tyr 1235 1240 1245
- Gly Gly Phe Ser Ser Thr Gln Asp Thr Val Val Ala Leu Asp Ala 1250 1255 1260
- Leu Ser Lys Tyr Gly Ala Ala Thr Phe Ser Lys Ser Gln Lys Thr 1265 1270 1275
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- Gln Val Asp Lys Ser Asn Arg Leu Leu Leu Gln Gln Val Ser Leu 1295 1300 1305
- Pro Tyr Ile Pro Gly Asn Tyr Thr Val Ser Val Ser Gly Glu Gly 1316 1320
- Cys Val Tyr Ala Gln Thr Thr Leu Arg Tyr Asn Val Pro Leu Glu 1325 1330 1335
- Lys Gln Gln Pro Ala Phe Ala Leu Lys Val Gln Thr Val Pro Leu 1349 1345 1350
- Thr Cys Asn Asn Pro Lys Gly Gln Asn Ser Phe Gln Ile Ser Leu 1355 1360 1365
- Glu Ile Ser Tyr Met Gly Ser Arg Pro Ala Ser Asn Met Val Ile 1370 1380
- Ala Asp Val Lys Met Leu Ser Gly Phe Ile Pro Leu Lys Pro Thr 1385 1390 1395
- Val Lys Lys Leu Glu Arg Leu Gly His Val Ser Arg Thr Glu Val 1400 1405 1410
- Thr Thr Asn Asn Val Leu Leu Tyr Leu Asp Gln Val Thr Asn Gln 1415 1420 1425
- Thr Leu Ser Phe Ser Phe Ile Ile Gln Gln Asp Ile Pro Val Lys 1430 1435 1440
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- Asp Glu Val Ala Phe Ala Glu Tyr Ser Ser Pro Cys Ser Ser Asp 1460 1465 1470
- Asp Gln Asn Val 1475